

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/590,122
Source: IFPP
Date Processed by STIC: 9/6/06

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/590,122

CRF Edit Date: 9/6/06
Edited by: KQ

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

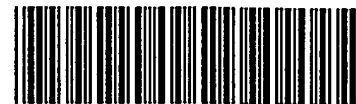
___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

___ Deleted: ☒ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

___ Moved responses to same line as heading/numeric identifier, specifically:

___ Other:



IFWP

RAW SEQUENCE LISTING

DATE: 09/06/2006

PATENT APPLICATION: US/10/590,122

TIME: 13:26:46

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\09052006\J590122.raw

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3 <110> APPLICANT: Yamakawa, Naomi
5 <120> TITLE OF INVENTION: DNA array for analyzing DNA methylation, process for
manufacturing the
6     same, and method for analyzing DNA methylation
8 <130> FILE REFERENCE: 2352.016
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/590,122
C--> 10 <141> CURRENT FILING DATE: 2006-08-18
10 <150> PRIOR APPLICATION NUMBER: PCT/JP2005/002490
11 <151> PRIOR FILING DATE: 2005-02-17
13 <160> NUMBER OF SEQ ID NOS: 4
15 <170> SOFTWARE: PatentIn version 3.1
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 20
19 <212> TYPE: DNA
20 <213> ORGANISM: Homo sapiens
22 <400> SEQUENCE: 1
23 attcgaaccc agtggaaatca                                20
26 <210> SEQ ID NO: 2
27 <211> LENGTH: 20
28 <212> TYPE: DNA
29 <213> ORGANISM: Homo sapiens
31 <400> SEQUENCE: 2
32 ctagcctgga gttgctaggg                                20
35 <210> SEQ ID NO: 3
36 <211> LENGTH: 20
37 <212> TYPE: DNA
38 <213> ORGANISM: Homo sapiens
40 <400> SEQUENCE: 3
41 gaactgcaaa gcacctgtga                                20
44 <210> SEQ ID NO: 4
45 <211> LENGTH: 19
46 <212> TYPE: DNA
47 <213> ORGANISM: Homo sapiens
49 <400> SEQUENCE: 4
50 gggctggagt ctgaactga                                19

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VERIFICATION SUMMARY

PATENT APPLICATION: **US/10/590,122**

DATE: 09/06/2006

TIME: 13:26:47

Input Set : **A:\PTO.KD.txt**

Output Set: **N:\CRF4\09052006\J590122.raw**

L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

Raw Sequence Listing before editing (for reference only)



IFWP

RAW SEQUENCE LISTING

DATE: 08/28/2006

PATENT APPLICATION: US/10/590,122

TIME: 09:52:57

Input Set : A:\PTO.RJ.txt

Output Set: N:\CRF4\08282006\J590122.raw

3 <110> APPLICANT: Yamakawa, Naomi
 5 <120> TITLE OF INVENTION: DNA array for analyzing DNA methylation, process for
 manufacturing the
 6 same, and method for analyzing DNA methylation
 8 <130> FILE REFERENCE: 2352.016
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/590,122
 C--> 10 <141> CURRENT FILING DATE: 2006-08-18
 10 <150> PRIOR APPLICATION NUMBER: PCT/JP2005/002490
 11 <151> PRIOR FILING DATE: 2005-02-17
 13 <160> NUMBER OF SEQ ID NOS: 4
 15 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
 Corrected Diskette Needed

(pg.1)

ERRORED SEQUENCES

44 <210> SEQ ID NO: 4
 45 <211> LENGTH: 19
 46 <212> TYPE: DNA
 47 <213> ORGANISM: Homo sapiens
 49 <400> SEQUENCE: 4
 50 gggctggagt ctgaactga
 E--> 57 1

19

deleted

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/590,122

DATE: 08/28/2006

TIME: 09:52:58

Input Set : A:\PTO.RJ.txt

Output Set: N:\CRF4\08282006\J590122.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:57 M:254 E: No. of Bases conflict, this line has no nucleotides.